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SEQUENCE LISTING

<110> COMPATIGÈNE INC.
PERREAULT, Claude
MCBRIDE, Kevin

<120> Mammalian SIMP, gene sequence and uses thereof in cancer therapy

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FOOTNOTES

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130				135				140							
Ala 145	Gly	Leu	Phe	Ala	Ala 150	Cys	Phe	Ile	Ala	Ile 155	Ser	Pro	Gly	Tyr	Thr 160
Ser	Arg	Ser	Val	Ala 165	Gly	Ser	Tyr	Asp	Asn 170	Glu	Gly	Ile	Ala	Ile 175	Phe
Ala	Leu	Gln	Phe 180	Thr	Tyr	Tyr	Leu	Trp 185	Val	Lys	Ser	Leu	Lys 190	Thr	Gly
Ser	Ile	Met 195	Trp	Ala	Ser	Leu	Cys 200	Ala	Leu	Ser	Tyr	Phe 205	Tyr	Met	Val
Ser	Ala 210	Trp	Gly	Gly	Tyr	Val 215	Phe	Ile	Ile	Asn	Leu 220	Ile	Pro	Leu	His
Ala 225	Leu	Ala	Leu	Ile	Ile 230	Met	Gly	Arg	Tyr	Ser 235	Ser	Arg	Leu	Phe	Val 240
Ser	Tyr	Thr	Ser	Phe 245	Tyr	Cys	Leu	Ala	Thr 250	Ile	Leu	Ser	Met	Gln 255	Val
Pro	Phe	Val	Gly 260	Phe	Gln	Pro	Val	Arg 265	Thr	Ser	Glu	His	Met 270	Pro	Ala
Phe	Gly	Val 275	Phe	Gly	Leu	Leu	Gln 280	Ile	Val	Ala	Leu	Met 285	His	Tyr	Ala
Arg	Asn 290	Arg	Ile	Thr	Arg	Gln 295	Gln	Phe	Met	Thr	Leu 300	Phe	Val	Gly	Gly
Leu 305	Thr	Ile	Leu	Gly	Ala 310	Leu	Ser	Val	Val	Val 315	Tyr	Phe	Ala	Leu	Val 320
Trp	Gly	Gly	Tyr	Val 325	Ala	Pro	Phe	Ser	Gly 330	Arg	Phe	Tyr	Ser	Leu 335	Trp
Asp	Thr	Gly	Tyr 340	Ala	Lys	Ile	His	Ile 345	Pro	Ile	Ile	Ala	Ser 350	Val	Ser
Glu	His	Gln 355	Pro	Thr	Thr	Trp	Val 360	Ser	Phe	Phe	Phe	Asp 365	Leu	His	Ile
Thr	Ala 370	Ala	Val	Phe	Pro	Val 375	Gly	Leu	Trp	Tyr	Cys 380	Ile	Lys	Lys	Val

Asn Asp Glu Arg Val Phe Ile Ile Leu Tyr Ala Val Ser Ala Val Tyr
385 390 395 400

Phe Ala Gly Val Met Val Arg Leu Met Leu Thr Leu Thr Pro Ala Val
405 410 415

Cys Val Leu Ala Gly Ile Gly Phe Ser Tyr Thr Phe Glu Lys Tyr Leu
420 425 430

Lys Asp Glu Glu Thr Lys Glu Arg Ser Ser Ser Gln Ser Gly Thr Thr
435 440 445

Lys Asp Glu Lys Leu Tyr Asp Lys Ala Ala Lys Asn Val Lys Ser Arg
450 455 460

Asn Ala Asn Asp Gly Asp Glu Ser Gly Val Ser Ser Asn Val Arg Thr
465 470 475 480

Ile Ile Ser Ile Ile Leu Val Ile Phe Leu Leu Met Phe Val Val His
485 490 495

Ala Thr Tyr Val Thr Ser Asn Ala Tyr Ser His Pro Ser Val Val Leu
500 505 510

Gln Ser Ser Thr Asn Asn Gly Asp Arg Ile Ile Met Asp Asp Phe Arg
515 520 525

Glu Ala Tyr His Trp Leu Arg Glu Asn Thr Ala Asp Asp Ala Arg Val
530 535 540

Met Ser Trp Trp Asp Tyr Gly Tyr Gln Ile Ala Gly Met Ala Asn Arg
545 550 555 560

Thr Thr Leu Val Asp Asn Asn Thr Trp Asn Asn Ser His Ile Ala Leu
565 570 575

Val Gly Lys Ala Met Ser Ser Asn Glu Ser Ala Ala Tyr Glu Ile Met
580 585 590

Thr Glu Leu Asp Val Asp Tyr Ile Leu Val Ile Phe Gly Gly Val Ile
595 600 605

Gly Tyr Ser Gly Asp Asp Ile Asn Lys Phe Leu Trp Met Val Arg Ile
610 615 620

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Thr Ala Ser Lys Ala
755